



SEQUENCE LISTING

<110> Institut Pasteur

<120> ANTIGÈNES DE PLASMODIUM FALCIPARUM ET LEURS APPLICATIONS
VACCINALES ET DIAGNOSTIQUES

<130> 000466-0035

<150> CA 2,345,206

<151> 2001-05-16

<150> CA 2,346,968

<151> 2001-05-23

<160> 18

<170> PatentIn version 3.1

<210> 1

<211> 192

<212> DNA

<213> Plasmodium falciparum

<400> 1

gaattccata tgcacgatta catatatgat gatcgatatct acaataatga taaagagaaa 60
aatgttataa aaagtataa taaaaatggt ataaaaagtg ataataaaaa tgattataaa 120
aagttaata aaaatgttat aaaaagtgat aataaaaatg ttataaaaag tgataataaa 180
aatgtggaat tc 192

<210> 2

<211> 351

<212> DNA

<213> Plasmodium falciparum

<400> 2

gaattcccag atcctcgaag taatgaccaa gaagatgcta ctgacgatgt ttagagaaaat 60
agtagagatg ataataatag tctctctaag agcgtagata atcaaagtaa tgttttaaat 120
agagaagatc ctattgcttc tgaaactgaa gttgtaagt aacctgagga ttcaagtagg 180
ataatgacta cagaagttcc aagtactact gtaaaacccc ctgatgaaaa acgatctgaa 240
gaagtaggag aaaaagaagc taaagaaatt aaagtagaac ctgttgtagc aagagccatt 300
ggagaaccaa tggaaaattc tgtgagcgta cagtcccctc ctaaggaatt c 351

<210> 3

<211> 64

<212> PRT

<213> Plasmodium falciparum

<400> 3

Glu Phe His Met His Asp Tyr Ile Tyr Asp Asp Arg Ile Tyr Asn Asn
1 5 10 15

Asp Lys Glu Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys
20 25 30

Ser Asp Asn Lys Asn Asp Tyr Lys Lys Cys Asn Lys Asn Val Ile Lys
35 40 45

Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Glu Phe
50 55 60

<210> 4

<211> 117

<212> PRT

<213> Plasmodium falciparum

<400> 4

Glu Phe Pro Asp Pro Arg Ser Asn Asp Gln Glu Asp Ala Thr Asp Asp
1 5 10 15

Val Val Glu Asn Ser Arg Asp Asp Asn Asn Ser Leu Ser Asn Ser Val
20 25 30

Asp Asn Gln Ser Asn Val Leu Asn Arg Glu Asp Pro Ile Ala Ser Glu
35 40 45

Thr Glu Val Val Ser Glu Pro Glu Asp Ser Ser Arg Ile Met Thr Thr
50 55 60

Glu Val Pro Ser Thr Thr Val Lys Pro Pro Asp Glu Lys Arg Ser Glu

65 70 75 80

Glu Val Gly Glu Lys Glu Ala Lys Glu Ile Lys Val Glu Pro Val Val
 85 90 95

Pro Arg Ala Ile Gly Glu Pro Met Glu Asn Ser Val Ser Val Gln Ser
 100 105 110

Pro Pro Lys Glu Phe
 115

<210> 5

<211> 5

<212> PRT

<213> Plasmodium falciparum

<400> 5

Asp Asn Lys Asn Val
1 5

<210> 6

<211> 5

<212> PRT

<213> Plasmodium falciparum

<400> 6

Asp Asn Lys Asn Asp
1 5

<210> 7

<211> 6

<212> PRT

<213> Plasmodium falciparum

<400> 7

Asp Lys Glu Lys Asn Val
1 5

<210> 8

<211> 7

<212> PRT

<213> Plasmodium falciparum

<400> 8

Lys Ser Asp Asn Lys Asn Val
1 5

<210> 9

<211> 1524

<212> DNA

<213> Plasmodium falciparum

<220>

<221> CDS

<222> (1)..(1524)

<223>

<400> 9

atg aag acg aca aaa gaa aat gac aat aat aac ata gta cat tat gta 48
Met Lys Thr Thr Lys Glu Asn Asp Asn Asn Asn Ile Val His Tyr Val
1 5 10 15

gat tgg ata aac cag att ttt aaa aag aat tct tta caa tgt gat tta 96
Asp Trp Ile Asn Gln Ile Phe Lys Lys Asn Ser Leu Gln Cys Asp Leu
20 25 30

tat ttt ttg gat gac aac aaa gaa aaa gat gtt agt aag aaa aga aaa 144
Tyr Phe Leu Asp Asp Asn Lys Glu Lys Asp Val Ser Lys Lys Arg Lys
35 40 45

gct caa ttg aag gat gaa tat gat aat ata tca agg agc aaa gaa aat 192
Ala Gln Leu Lys Asp Glu Tyr Asp Asn Ile Ser Arg Ser Lys Glu Asn
50 55 60

att aat aat tcc aaa aaa ata aaa aat gaa tta agt ata aaa gat aat 240
Ile Asn Asn Ser Lys Lys Ile Lys Asn Glu Leu Ser Ile Lys Asp Asn
65 70 75 80

atg cac gat tac ata tat gat gat cgt atc tac aat aat gat aaa gag 288
Met His Asp Tyr Ile Tyr Asp Asp Arg Ile Tyr Asn Asn Asp Lys Glu
85 90 95

aaa aat gtt ata aaa agt gat aat aaa aat gtt ata aaa agt gat aat 336
Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn
100 105 110

aaa aat gat tat aaa aag tgt aat aaa aat gtt ata aaa agt gat aat 384
Lys Asn Asp Tyr Lys Lys Cys Asn Lys Asn Val Ile Lys Ser Asp Asn
115 120 125

aaa aat gtt ata aaa agt gat aat aaa aat gtt ata aaa agt gat aat 432
Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn
130 135 140

aaa aat gtt ata aaa agt gat tat aaa agt gat gat aga aat gct tgt 480
Lys Asn Val Ile Lys Ser Asp Tyr Lys Ser Asp Asp Arg Asn Ala Cys

145	150	155	160	
gat att tat aaa agt aat aaa aaa aat gtt cct gat aat tgc cat ata				528
Asp Ile Tyr Lys Ser Asn Lys Lys Asn Val Pro Asp Asn Cys His Ile				
165	170	175		
tat gat gat aat agt tca gtt gaa aat tta gat gga aaa aat aaa tta				576
Tyr Asp Asp Asn Ser Ser Val Glu Asn Leu Asp Gly Lys Asn Lys Leu				
180	185	190		
aat aat ata agg aac ata cat aat gat aac tca tct tca tgc gat ata				624
Asn Asn Ile Arg Asn Ile His Asn Asp Asn Ser Ser Ser Cys Asp Ile				
195	200	205		
tcc gat ata aaa agt gaa gat gaa tat ata gaa cca tat gaa aaa aag				672
Ser Asp Ile Lys Ser Glu Asp Glu Tyr Ile Glu Pro Tyr Glu Lys Lys				
210	215	220		
aat gaa gaa aat ata aat gaa tat aag aat aag aaa aat ata gcc aat				720
Asn Glu Glu Asn Ile Asn Glu Tyr Lys Asn Lys Lys Asn Ile Ala Asn				
225	230	235	240	
gaa aat ata aaa gaa gga aag agt tca att tat aat gat gaa cat aat				768
Glu Asn Ile Lys Glu Gly Lys Ser Ser Ile Tyr Asn Asp Glu His Asn				
245	250	255		
tat aat tca tta tta tat aat tct tgt aat ggt gaa ata agt aag atc				816
Tyr Asn Ser Leu Leu Tyr Asn Ser Cys Asn Gly Glu Ile Ser Lys Ile				
260	265	270		
aac aaa ata agt agt cat aat aat att gat aat aat atg gat aat tat				864
Asn Lys Ile Ser Ser His Asn Asn Ile Asp Asn Asn Met Asp Asn Tyr				
275	280	285		
aat acg ttt gca aat gtg aat aat ttt ata ata tat tcc tca gat gat				912
Asn Thr Phe Ala Asn Val Asn Asn Phe Ile Ile Tyr Ser Ser Asp Asp				
290	295	300		
gaa gat aat ata tca aat tat tat aat ggt aaa gac gta tta aat gat				960
Glu Asp Asn Ile Ser Asn Tyr Tyr Asn Gly Lys Asp Val Leu Asn Asp				
305	310	315	320	
gag att atg ttc cct ata aaa ttt aat ttt gaa aaa tta aaa aaa aat				1008
Glu Ile Met Phe Pro Ile Lys Phe Asn Phe Glu Lys Leu Lys Lys Asn				
325	330	335		

att tat gta ata gag cat ata gac aaa ata tat tat gat aca ttt tta 1056
 Ile Tyr Val Ile Glu His Ile Asp Lys Ile Tyr Tyr Asp Thr Phe Leu
 340 345 350

aat aaa aat cca agt gaa aaa agt gtt ttt atg aat gat gaa tct act 1104
 Asn Lys Asn Pro Ser Glu Lys Ser Val Phe Met Asn Asp Glu Ser Thr
 355 360 365

ggc tat ttg aaa aat gat gtg aat gac aaa tgt gtt gtt gat aat ata 1152
 Gly Tyr Leu Lys Asn Asp Val Asn Asp Lys Cys Val Val Asp Asn Ile
 370 375 380

aat gtt att aat cct tct agt gtg aat acg ttg agt aat att tca aat 1200
 Asn Val Ile Asn Pro Ser Ser Val Asn Thr Leu Ser Asn Ile Ser Asn
 385 390 395 400

att agg aat gaa aaa ata gaa aat aat aat aag aat gaa aaa tta ata 1248
 Ile Arg Asn Glu Lys Ile Glu Asn Asn Asn Lys Asn Glu Lys Leu Ile
 405 410 415

aaa tca tat cct aca caa tca aaa aat gtt atg agt aca ttt tcc ttt 1296
 Lys Ser Tyr Pro Thr Gln Ser Lys Asn Val Met Ser Thr Phe Ser Phe
 420 425 430

tgg aat att gaa aag gag aca ttt ata aca aaa cct ttg tat gca caa 1344
 Trp Asn Ile Glu Lys Glu Thr Phe Ile Thr Lys Pro Leu Tyr Ala Gln
 435 440 445

aat ttg aga aaa aaa caa ttt agt tta tta gat gaa tct gaa gag atg 1392
 Asn Leu Arg Lys Lys Gln Phe Ser Leu Leu Asp Glu Ser Glu Glu Met
 450 455 460

ata aga aat tat tca tct aat caa tat tct ata aaa ttt gta cca aga 1440
 Ile Arg Asn Tyr Ser Ser Asn Gln Tyr Ser Ile Lys Phe Val Pro Arg
 465 470 475 480

cat tta tta tat gta atg agt caa gtt gct tct cga tcc ttt ttt gat 1488
 His Leu Leu Tyr Val Met Ser Gln Val Ala Ser Arg Ser Phe Phe Asp
 485 490 495

cct tta tat aga aag cag tta ttt ttt cgt tac taa 1524
 Pro Leu Tyr Arg Lys Gln Leu Phe Phe Arg Tyr
 500 505

<210> 10

<211> 507

<212> PRT

<213> Plasmodium falciparum

<400> 10

Met Lys Thr Thr Lys Glu Asn Asp Asn Asn Ile Val His Tyr Val
1 5 10 15

Asp Trp Ile Asn Gln Ile Phe Lys Lys Asn Ser Leu Gln Cys Asp Leu
20 25 30

Tyr Phe Leu Asp Asp Asn Lys Glu Lys Asp Val Ser Lys Lys Arg Lys
35 40 45

Ala Gln Leu Lys Asp Glu Tyr Asp Asn Ile Ser Arg Ser Lys Glu Asn
50 55 60

Ile Asn Asn Ser Lys Lys Ile Lys Asn Glu Leu Ser Ile Lys Asp Asn
65 70 75 80

Met His Asp Tyr Ile Tyr Asp Asp Arg Ile Tyr Asn Asn Asp Lys Glu
85 90 95

Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn
100 105 110

Lys Asn Asp Tyr Lys Lys Cys Asn Lys Asn Val Ile Lys Ser Asp Asn
115 120 125

Lys Asn Val Ile Lys Ser Asp Asn Lys Asn Val Ile Lys Ser Asp Asn
130 135 140

Lys Asn Val Ile Lys Ser Asp Tyr Lys Ser Asp Asp Arg Asn Ala Cys
145 150 155 160

Asp Ile Tyr Lys Ser Asn Lys Lys Asn Val Pro Asp Asn Cys His Ile
165 170 175

Tyr Asp Asp Asn Ser Ser Val Glu Asn Leu Asp Gly Lys Asn Lys Leu
180 185 190

Asn Asn Ile Arg Asn Ile His Asn Asp Asn Ser Ser Ser Cys Asp Ile
195 200 205

Ser Asp Ile Lys Ser Glu Asp Glu Tyr Ile Glu Pro Tyr Glu Lys Lys
210 215 220

Asn Glu Glu Asn Ile Asn Glu Tyr Lys Asn Lys Lys Asn Ile Ala Asn
225 230 235 240

Glu Asn Ile Lys Glu Gly Lys Ser Ser Ile Tyr Asn Asp Glu His Asn
245 250 255

Tyr Asn Ser Leu Leu Tyr Asn Ser Cys Asn Gly Glu Ile Ser Lys Ile
260 265 270

Asn Lys Ile Ser Ser His Asn Asn Ile Asp Asn Asn Met Asp Asn Tyr
275 280 285

Asn Thr Phe Ala Asn Val Asn Asn Phe Ile Ile Tyr Ser Ser Asp Asp
290 295 300

Glu Asp Asn Ile Ser Asn Tyr Tyr Asn Gly Lys Asp Val Leu Asn Asp
305 310 315 320

Glu Ile Met Phe Pro Ile Lys Phe Asn Phe Glu Lys Leu Lys Lys Asn

325 330 335

Ile Tyr Val Ile Glu His Ile Asp Lys Ile Tyr Tyr Asp Thr Phe Leu
340 345 350

Asn Lys Asn Pro Ser Glu Lys Ser Val Phe Met Asn Asp Glu Ser Thr
355 360 365

Gly Tyr Leu Lys Asn Asp Val Asn Asp Lys Cys Val Val Asp Asn Ile
370 375 380

Asn Val Ile Asn Pro Ser Ser Val Asn Thr Leu Ser Asn Ile Ser Asn
385 390 395 400

Ile Arg Asn Glu Lys Ile Glu Asn Asn Asn Lys Asn Glu Lys Leu Ile
405 410 415

Lys Ser Tyr Pro Thr Gln Ser Lys Asn Val Met Ser Thr Phe Ser Phe
420 425 430

Trp Asn Ile Glu Lys Glu Thr Phe Ile Thr Lys Pro Leu Tyr Ala Gln
435 440 445

Asn Leu Arg Lys Lys Gln Phe Ser Leu Leu Asp Glu Ser Glu Glu Met
450 455 460

Ile Arg Asn Tyr Ser Ser Asn Gln Tyr Ser Ile Lys Phe Val Pro Arg
465 470 475 480

His Leu Leu Tyr Val Met Ser Gln Val Ala Ser Arg Ser Phe Phe Asp
485 490 495

Pro Leu Tyr Arg Lys Gln Leu Phe Phe Arg Tyr
500 505

<210> 11

<211> 5050

<212> DNA

<213> Plasmodium falciparum

<220>

<221> CDS

<222> (1)..(4464)

<223>

<220>

<221> CDS

<222> (4557)..(4634)

<223>

<220>

<221> CDS

<222> (4751)..(4837)

<223>

<220>

<221> CDS

<222> (4973)..(5047)

<223>

<400> 11

atg aaa ggg aaa atg aat atg tgt ttg ttt ttt ttc tat tct ata tta 48
Met Lys Gly Lys Met Asn Met Cys Leu Phe Phe Phe Tyr Ser Ile Leu
1 5 10 15

tat gtt gta tta tgt acc tat gta tta ggt ata agt gaa gag tat ttg 96
Tyr Val Val Leu Cys Thr Tyr Val Leu Gly Ile Ser Glu Glu Tyr Leu
20 25 30

aag gaa agg ccc caa ggt tta aat gtt gag act aat aat aat aat aat 144
Lys Glu Arg Pro Gln Gly Leu Asn Val Glu Thr Asn Asn Asn Asn Asn
35 40 45

aat aat aat aat aat aat agt aat agt aac gat gcg atg tct ttt gta 192
Asn Asn Asn Asn Asn Asn Ser Asn Ser Asn Asp Ala Met Ser Phe Val
50 55 60

aat gaa gta ata agg ttt ata gaa aac gag aag gat gat aaa gaa gat 240
Asn Glu Val Ile Arg Phe Ile Glu Asn Glu Lys Asp Asp Lys Glu Asp
65 70 75 80

aaa aaa gtg aag ata ata tct aga cct gtt gag aat aca tta cat aga 288
Lys Lys Val Lys Ile Ile Ser Arg Pro Val Glu Asn Thr Leu His Arg
85 90 95

tat cca gtt agt tct ttt ctg aat atc aaa aag tat ggt agg aaa ggg 336
Tyr Pro Val Ser Ser Phe Leu Asn Ile Lys Lys Tyr Gly Arg Lys Gly
100 105 110

gaa tat ttg aat aga aat agt ttt gtt caa aga tca tat ata agg ggt 384
Glu Tyr Leu Asn Arg Asn Ser Phe Val Gln Arg Ser Tyr Ile Arg Gly
115 120 125

tgt aaa gga aaa aga agc aca cat aca tgg ata tgt gaa aat aaa ggg 432
Cys Lys Gly Lys Arg Ser Thr His Thr Trp Ile Cys Glu Asn Lys Gly
130 135 140

aat aat aat ata tgt att cct gat aga cgt gta caa tta tgt ata aca 480
Asn Asn Asn Ile Cys Ile Pro Asp Arg Arg Val Gln Leu Cys Ile Thr
145 150 155 160

gct ctt caa gat tta aaa aat tca gga tct gaa acg act gat aga aaa 528
 Ala Leu Gln Asp Leu Lys Asn Ser Gly Ser Glu Thr Thr Asp Arg Lys
 165 170 175

tta tta aga gat aaa gta ttt gat tca gct atg tat gaa act gat ttg 576
 Leu Leu Arg Asp Lys Val Phe Asp Ser Ala Met Tyr Glu Thr Asp Leu
 180 185 190

tta tgg aat aaa tat ggt ttt cgt gga ttt gat gat ttt tgt gac gat 624
 Leu Trp Asn Lys Tyr Gly Phe Arg Gly Phe Asp Asp Phe Cys Asp Asp
 195 200 205

gta aaa aat agt tat tta gat tat aaa gat gtt ata ttt gga acc gat 672
 Val Lys Asn Ser Tyr Leu Asp Tyr Lys Asp Val Ile Phe Gly Thr Asp
 210 215 220

tta gat aaa aat aat ata tca aag tta gta gag gaa tca tta aaa cgt 720
 Leu Asp Lys Asn Asn Ile Ser Lys Leu Val Glu Glu Ser Leu Lys Arg
 225 230 235 240

ttt ttt aaa aaa gat agt agt gta ctt aat cct act gct tgg tgg aga 768
 Phe Phe Lys Lys Asp Ser Ser Val Leu Asn Pro Thr Ala Trp Trp Arg
 245 250 255

agg tat gga aca aga cta tgg aaa act atg ata cag cca tat gct cat 816
 Arg Tyr Gly Thr Arg Leu Trp Lys Thr Met Ile Gln Pro Tyr Ala His
 260 265 270

tta gga tgt aga aaa cct gat gag aat gaa cct cag ata aat aga tgg 864
 Leu Gly Cys Arg Lys Pro Asp Glu Asn Glu Pro Gln Ile Asn Arg Trp
 275 280 285

att ctg gaa tgg ggg aaa tat aat tgt aga tta atg aag gag aaa gaa 912
 Ile Leu Glu Trp Gly Lys Tyr Asn Cys Arg Leu Met Lys Glu Lys Glu
 290 295 300

aaa ttg tta aca gga gaa tgt tct gtt aat aga aaa aaa tct gac tgc 960
 Lys Leu Leu Thr Gly Glu Cys Ser Val Asn Arg Lys Lys Ser Asp Cys
 305 310 315 320

tca acc gga tgt aat aat gag tgt tat acc tat agg agt ctt att aat 1008
 Ser Thr Gly Cys Asn Asn Glu Cys Tyr Thr Tyr Arg Ser Leu Ile Asn
 325 330 335

aga caa aga tat gag gtc tct ata tta gga aaa aaa tat att aaa gta 1056
 Arg Gln Arg Tyr Glu Val Ser Ile Leu Gly Lys Lys Tyr Ile Lys Val

340	345	350	
gta cga tat act ata ttt agg aga aaa ata gtt caa cct gat aat gct 1104			
Val Arg Tyr Thr Ile Phe Arg Arg Lys Ile Val Gln Pro Asp Asn Ala			
355	360	365	
ttg gat ttt tta aaa tta aat tgt tct gag tgt aag gat att gat ttt 1152			
Leu Asp Phe Leu Lys Leu Asn Cys Ser Glu Cys Lys Asp Ile Asp Phe			
370	375	380	
aaa ccc ttt ttt gaa ttt gaa tat ggt aaa tat gaa gaa aaa tgt atg 1200			
Lys Pro Phe Phe Glu Phe Glu Tyr Gly Lys Tyr Glu Glu Lys Cys Met			
385	390	395	400
tgt caa tca tat att gat tta aaa atc caa ttt aaa aat aat gat att 1248			
Cys Gln Ser Tyr Ile Asp Leu Lys Ile Gln Phe Lys Asn Asn Asp Ile			
405	410	415	
tgt tca ttt aat gct caa aca gat act gtt tct agc gat aaa aga ttt 1296			
Cys Ser Phe Asn Ala Gln Thr Asp Thr Val Ser Ser Asp Lys Arg Phe			
420	425	430	
tgt ctt gaa aag aaa gaa ttt aaa cca tgg aaa tgt gat aaa aat tct 1344			
Cys Leu Glu Lys Lys Glu Phe Lys Pro Trp Lys Cys Asp Lys Asn Ser			
435	440	445	
ttt gaa aca gtt cat cat aaa ggt gta tgt gtg tca ccg aga aga caa 1392			
Phe Glu Thr Val His His Lys Gly Val Cys Val Ser Pro Arg Arg Gln			
450	455	460	
ggc ttt tgt tta gga aat ttg aac tat cta ctg aat gat gat att tat 1440			
Gly Phe Cys Leu Gly Asn Leu Asn Tyr Leu Leu Asn Asp Asp Ile Tyr			
465	470	475	480
aat gta cat aat tca caa cta ctt atc gaa att ata atg gct tct aaa 1488			
Asn Val His Asn Ser Gln Leu Leu Ile Glu Ile Ile Met Ala Ser Lys			
485	490	495	
caa gaa gga aag tta tta tgg aaa aaa cat gga aca ata ctt gat aac 1536			
Gln Glu Gly Lys Leu Leu Trp Lys Lys His Gly Thr Ile Leu Asp Asn			
500	505	510	
cag aat gca tgc aaa tat ata aat gat agt tat gtt gat tat aaa gat 1584			
Gln Asn Ala Cys Lys Tyr Ile Asn Asp Ser Tyr Val Asp Tyr Lys Asp			
515	520	525	

ata gtt att gga aat gat tta tgg aat gat aac aac tct ata aaa gtt 1632
 Ile Val Ile Gly Asn Asp Leu Trp Asn Asp Asn Asn Ser Ile Lys Val
 530 535 540

caa aat aat tta aat tta att ttt gaa aga aat ttt ggt tat aaa gtt 1680
 Gln Asn Asn Leu Asn Leu Ile Phe Glu Arg Asn Phe Gly Tyr Lys Val
 545 550 555 560

gga aga aat aaa ctc ttt aaa aca att aaa gaa tta aaa aat gta tgg 1728
 Gly Arg Asn Lys Leu Phe Lys Thr Ile Lys Glu Leu Lys Asn Val Trp
 565 570 575

tgg ata tta aat aga aat aaa gta tgg gaa tca atg aga tgt gga att 1776
 Trp Ile Leu Asn Arg Asn Lys Val Trp Glu Ser Met Arg Cys Gly Ile
 580 585 590

gac gaa gta gat caa cgt aga aaa act tgt gaa aga ata gat gaa cta 1824
 Asp Glu Val Asp Gln Arg Arg Lys Thr Cys Glu Arg Ile Asp Glu Leu
 595 600 605

gaa aac atg cca caa ttc ttt aga tgg ttt tca caa tgg gca cat ttc 1872
 Glu Asn Met Pro Gln Phe Phe Arg Trp Phe Ser Gln Trp Ala His Phe
 610 615 620

ttt tgt aag gaa aaa gaa tat tgg gaa tta aaa tta aat gat aaa tgt 1920
 Phe Cys Lys Glu Lys Glu Tyr Trp Glu Leu Lys Leu Asn Asp Lys Cys
 625 630 635 640

aca ggt aat aat gga aaa tcc tta tgt cag gat aaa aca tgt caa aat 1968
 Thr Gly Asn Asn Gly Lys Ser Leu Cys Gln Asp Lys Thr Cys Gln Asn
 645 650 655

gtg tgt act aat atg aat tat tgg aca tat act aga aaa tta gct tat 2016
 Val Cys Thr Asn Met Asn Tyr Trp Thr Tyr Thr Arg Lys Leu Ala Tyr
 660 665 670

gaa ata caa tcc gta aaa tat gat aaa gat aga aaa tta ttt agt ctt 2064
 Glu Ile Gln Ser Val Lys Tyr Asp Lys Asp Arg Lys Leu Phe Ser Leu
 675 680 685

gct aaa gac aaa aat gta act aca ttt tta aag gaa aat gca aaa aat 2112
 Ala Lys Asp Lys Asn Val Thr Thr Phe Leu Lys Glu Asn Ala Lys Asn
 690 695 700

tgt tct aat ata gat ttt aca aaa ata ttc gat cag ctt gac aaa ctc 2160
 Cys Ser Asn Ile Asp Phe Thr Lys Ile Phe Asp Gln Leu Asp Lys Leu

705	710	715	720	
ttt aag gaa aga tgt tca tgt atg gat aca caa gtt tta gaa gta aaa				2208
Phe Lys Glu Arg Cys Ser Cys Met Asp Thr Gln Val Leu Glu Val Lys				
725	730	735		
aac aaa gaa atg tta tct ata gac tca aat agt gaa gat gcg aca gat				2256
Asn Lys Glu Met Leu Ser Ile Asp Ser Asn Ser Glu Asp Ala Thr Asp				
740	745	750		
ata agt gag aaa aat gga gag gaa gaa tta tat gta aat cac aat tct				2304
Ile Ser Glu Lys Asn Gly Glu Glu Glu Leu Tyr Val Asn His Asn Ser				
755	760	765		
gtg agt gtc gca agt ggt aat aaa gaa atc gaa aag agt aag gat gaa				2352
Val Ser Val Ala Ser Gly Asn Lys Glu Ile Glu Lys Ser Lys Asp Glu				
770	775	780		
aag caa cct gaa aaa gaa gca aaa caa act aat gga act tta acc gta				2400
Lys Gln Pro Glu Lys Glu Ala Lys Gln Thr Asn Gly Thr Leu Thr Val				
785	790	795	800	
cga act gac aaa gat tca gat aga aac aaa gga aaa gat aca gct act				2448
Arg Thr Asp Lys Asp Ser Asp Arg Asn Lys Gly Lys Asp Thr Ala Thr				
805	810	815		
gat aca aaa aat tca cct gaa aat tta aaa gta cag gaa cat gga aca				2496
Asp Thr Lys Asn Ser Pro Glu Asn Leu Lys Val Gln Glu His Gly Thr				
820	825	830		
aat gga gaa aca ata aaa gaa gaa cca cca aaa tta cct gaa tca tct				2544
Asn Gly Glu Thr Ile Lys Glu Glu Pro Pro Lys Leu Pro Glu Ser Ser				
835	840	845		
gaa aca tta caa tca caa gaa caa tta gaa gca gaa gca caa aaa caa				2592
Glu Thr Leu Gln Ser Gln Glu Gln Leu Glu Ala Glu Ala Gln Lys Gln				
850	855	860		
aaa caa gaa gaa gaa cca aaa aaa aaa caa gaa gaa gaa cca aaa aaa				2640
Lys Gln Glu Glu Glu Pro Lys Lys Lys Gln Glu Glu Glu Pro Lys Lys				
865	870	875	880	
aaa caa gaa gaa gaa caa aaa cga gaa caa gaa caa aaa caa gaa caa				2688
Lys Gln Glu Glu Glu Gln Lys Arg Glu Gln Glu Gln Lys Gln Glu Gln				
885	890	895		

gaa gaa gaa gaa caa aaa caa gaa gaa gaa caa caa ata caa gat caa 2736
 Glu Glu Glu Glu Gln Lys Gln Glu Glu Glu Gln Gln Ile Gln Asp Gln
 900 905 910

tca caa agt gga tta gat caa tcc tca aaa gta gga gta gcg agt gaa 2784
 Ser Gln Ser Gly Leu Asp Gln Ser Ser Lys Val Gly Val Ala Ser Glu
 915 920 925

caa aat gaa att tct tca gga caa gaa caa aac gta aaa agc tct tca 2832
 Gln Asn Glu Ile Ser Ser Gly Gln Glu Gln Asn Val Lys Ser Ser Ser
 930 935 940

cct gaa gta gtt cca caa gaa aca act agt gaa aat ggg tca tca caa 2880
 Pro Glu Val Val Pro Gln Glu Thr Thr Ser Glu Asn Gly Ser Ser Gln
 945 950 955 960

gac aca aaa ata tca agt act gaa cca aat gag aat tct gtt gta gat 2928
 Asp Thr Lys Ile Ser Ser Thr Glu Pro Asn Glu Asn Ser Val Val Asp
 965 970 975

aga gca aca gat agt atg aat tta gat cct gaa aag gtt cat aat gaa 2976
 Arg Ala Thr Asp Ser Met Asn Leu Asp Pro Glu Lys Val His Asn Glu
 980 985 990

aat atg agt gat cca aat aca aat act gaa cca gat gca tct tta aaa 3024
 Asn Met Ser Asp Pro Asn Thr Asn Thr Glu Pro Asp Ala Ser Leu Lys
 995 1000 1005

gat gat aag aag gaa gtt gat gat gcc aaa aaa gaa ctt caa tct 3069
 Asp Asp Lys Lys Glu Val Asp Asp Ala Lys Lys Glu Leu Gln Ser
 1010 1015 1020

act gta tca aga att gaa tct aat gaa cag gac gtt caa agt aca 3114
 Thr Val Ser Arg Ile Glu Ser Asn Glu Gln Asp Val Gln Ser Thr
 1025 1030 1035

cca ccc gaa gat act cct act gtt gaa gga aaa gta gga gat aaa 3159
 Pro Pro Glu Asp Thr Pro Thr Val Glu Gly Lys Val Gly Asp Lys
 1040 1045 1050

gca gaa atg tta act tct ccg cat gcg aca gat aat tct gag tcg 3204
 Ala Glu Met Leu Thr Ser Pro His Ala Thr Asp Asn Ser Glu Ser
 1055 1060 1065

gaa tca ggt tta aat cca act gat gac att aaa aca act gat ggt 3249
 Glu Ser Gly Leu Asn Pro Thr Asp Asp Ile Lys Thr Thr Asp Gly

1070	1075	1080	
gtt gtt aaa gaa caa gaa ata tta ggg gga ggt gaa agt gca act	3294		
Val Val Lys Glu Gln Glu Ile Leu Gly Gly Gly Glu Ser Ala Thr			
1085	1090	1095	
gaa aca tca aaa agt aat tta gaa aaa cct aag gat gtt gaa cct	3339		
Glu Thr Ser Lys Ser Asn Leu Glu Lys Pro Lys Asp Val Glu Pro			
1100	1105	1110	
tct cat gaa ata tct gaa cct gtt ctt tct ggt aca act ggt aaa	3384		
Ser His Glu Ile Ser Glu Pro Val Leu Ser Gly Thr Thr Gly Lys			
1115	1120	1125	
gaa gaa tca gag tta tta aaa agt aaa tcg ata gag acg aag ggg	3429		
Glu Glu Ser Glu Leu Leu Lys Ser Lys Ser Ile Glu Thr Lys Gly			
1130	1135	1140	
gaa aca gat cct cga agt aat gac caa gaa gat gct act gac gat	3474		
Glu Thr Asp Pro Arg Ser Asn Asp Gln Glu Asp Ala Thr Asp Asp			
1145	1150	1155	
gtt gta gaa aat agt aga gat gat aat aat agt ctc tct aat agc	3519		
Val Val Glu Asn Ser Arg Asp Asp Asn Asn Ser Leu Ser Asn Ser			
1160	1165	1170	
gta gat aat caa agt aat gtt tta aat aga gaa gat cct att gct	3564		
Val Asp Asn Gln Ser Asn Val Leu Asn Arg Glu Asp Pro Ile Ala			
1175	1180	1185	
tct gaa act gaa gtt gta agt gaa cct gag gat tca agt agg ata	3609		
Ser Glu Thr Glu Val Val Ser Glu Pro Glu Asp Ser Ser Arg Ile			
1190	1195	1200	
atg act aca gaa gtt cca agt act act gta aaa ccc cct gat gaa	3654		
Met Thr Thr Glu Val Pro Ser Thr Thr Val Lys Pro Pro Asp Glu			
1205	1210	1215	
aaa cga tct gaa gaa gta gga gaa aaa gaa gct aaa gaa att aaa	3699		
Lys Arg Ser Glu Glu Val Gly Glu Lys Glu Ala Lys Glu Ile Lys			
1220	1225	1230	
gta gaa cct gtt gta cca aga gcc att gga gaa cca atg gaa aat	3744		
Val Glu Pro Val Val Pro Arg Ala Ile Gly Glu Pro Met Glu Asn			
1235	1240	1245	

tct gtg agc gta cag tcc cct cct aat gta gaa gat gtt gaa aaa 3789
 Ser Val Ser Val Gln Ser Pro Pro Asn Val Glu Asp Val Glu Lys
 1250 1255 1260

gaa aca ttg ata tct gag aat aat gga tta cat aat gat aca cac 3834
 Glu Thr Leu Ile Ser Glu Asn Asn Gly Leu His Asn Asp Thr His
 1265 1270 1275

aga gga aat atc agt gaa aag gat tta atc gat att cat ttg tta 3879
 Arg Gly Asn Ile Ser Glu Lys Asp Leu Ile Asp Ile His Leu Leu
 1280 1285 1290

aga aat gaa gcg ggt agt aca ata tta gat gat tct aga aga aat 3924
 Arg Asn Glu Ala Gly Ser Thr Ile Leu Asp Asp Ser Arg Arg Asn
 1295 1300 1305

gga gaa atg aca gaa ggt agc gaa agt gat gtt gga gaa tta caa 3969
 Gly Glu Met Thr Glu Gly Ser Glu Ser Asp Val Gly Glu Leu Gln
 1310 1315 1320

gaa cat aat ttt agc aca caa caa aaa gat gaa aaa gat ttt gac 4014
 Glu His Asn Phe Ser Thr Gln Gln Lys Asp Glu Lys Asp Phe Asp
 1325 1330 1335

caa att gcg agc gat aga gaa aaa gaa gaa att caa aaa tta ctt 4059
 Gln Ile Ala Ser Asp Arg Glu Lys Glu Glu Ile Gln Lys Leu Leu
 1340 1345 1350

aat ata gga cat gaa gag gat gaa gat gta tta aaa atg gat aga 4104
 Asn Ile Gly His Glu Glu Asp Glu Asp Val Leu Lys Met Asp Arg
 1355 1360 1365

aca gag gat agt atg agt gat gga gtt aat agt cat ttg tat tat 4149
 Thr Glu Asp Ser Met Ser Asp Gly Val Asn Ser His Leu Tyr Tyr
 1370 1375 1380

aat aat cta tca agt gaa gaa aaa atg gaa caa tat aat aat aga 4194
 Asn Asn Leu Ser Ser Glu Glu Lys Met Glu Gln Tyr Asn Asn Arg
 1385 1390 1395

gat gct tct aaa gat aga gaa gaa ata ttg aat agg tca aac aca 4239
 Asp Ala Ser Lys Asp Arg Glu Glu Ile Leu Asn Arg Ser Asn Thr
 1400 1405 1410

aat aca tgt tct aat gaa cat tca tta aaa tat tgt caa tat atg 4284
 Asn Thr Cys Ser Asn Glu His Ser Leu Lys Tyr Cys Gln Tyr Met

1415	1420	1425	
gaa aga aat aag gat tta tta gaa aca tgt tct gaa gac aaa agg 4329			
Glu Arg	Asn Lys Asp Leu Leu	Glu Thr Cys Ser Glu	Asp Lys Arg
1430	1435	1440	
tta cat tta tgt tgt gaa ata tca gat tat tgt tta aaa ttt ttc 4374			
Leu His	Leu Cys Cys Glu Ile	Ser Asp Tyr Cys Leu	Lys Phe Phe
1445	1450	1455	
aat cct aaa tcg ata gaa tac ttt gat tgt aca caa aaa gaa ttt 4419			
Asn Pro	Lys Ser Ile Glu Tyr	Phe Asp Cys Thr Gln	Lys Glu Phe
1460	1465	1470	
gat gac cct aca tat aat tgt ttt aga aaa caa aga ttt aca agt 4464			
Asp Asp	Pro Thr Tyr Asn Cys	Phe Arg Lys Gln Arg	Phe Thr Ser
1475	1480	1485	
atgcatgtt ataaaattaa aaacaatata cattaatatg ttaataaaaa aaataatata 4524			
ttttttctc tttttcttt ttttaatat gt atg cat tat att gcc ggg ggt 4577			
	Met His	Tyr Ile Ala Gly Gly	
	1490	1495	
ggg ata ata gcc ctt tta ttg ttt att tta ggt tca gcc agc tat 4622			
Gly Ile Ile	Ala Leu Leu Leu Phe Ile Leu	Gly Ser Ala Ser Tyr	
1500	1505	1510	
agg aag aat ttg taagaaaaaa aggatgaaga aatataaaca aaaatataaa 4674			
Arg Lys Asn Leu			
tatatgcata tatatttaag tattataaga acatatatat aaataaatat gtatatTTTT 4734			
attttattat tatagg gat gat gaa aaa gga ttc tac gat tct aat tta 4783			
Asp Asp	Glu Lys Gly Phe Tyr Asp	Ser Asn Leu	
1515	1520	1525	
aat gat tct gct ttt gaa tat aat aat aat aaa tat aat aaa tta 4828			
Asn Asp	Ser Ala Phe Glu Tyr Asn Asn Asn	Lys Tyr Asn Lys Leu	
1530	1535	1540	
cct tat atg tgtaaggaaa aaactaaaaa acaaaaaaaa aaaaaaatat 4877			
Pro Tyr Met			

atatatatat atatataattt acggatgcat ttccacattc ctattatttc ttattcttat 4937

aatttttatt atttatttat ttattttttt ttttc gta gtt gat caa caa ata 4990
Val Val Asp Gln Gln Ile
1545

aat gta gta aat tct gat tta tat tcg gag ggt att tat gat gac 5035
Asn Val Val Asn Ser Asp Leu Tyr Ser Glu Gly Ile Tyr Asp Asp
1550 1555 1560

aca acg aca ttt taa 5050
Thr Thr Thr Phe
1565

<210> 12

<211> 1568

<212> PRT

<213> Plasmodium falciparum

<400> 12

Met Lys Gly Lys Met Asn Met Cys Leu Phe Phe Phe Tyr Ser Ile Leu
1 5 10 15

Tyr Val Val Leu Cys Thr Tyr Val Leu Gly Ile Ser Glu Glu Tyr Leu
20 25 30

Lys Glu Arg Pro Gln Gly Leu Asn Val Glu Thr Asn Asn Asn Asn Asn
35 40 45

Asn Asn Asn Asn Asn Asn Ser Asn Ser Asn Asp Ala Met Ser Phe Val
50 55 60

Asn Glu Val Ile Arg Phe Ile Glu Asn Glu Lys Asp Asp Lys Glu Asp
65 70 75 80

Lys Lys Val Lys Ile Ile Ser Arg Pro Val Glu Asn Thr Leu His Arg
85 90 95

Tyr Pro Val Ser Ser Phe Leu Asn Ile Lys Lys Tyr Gly Arg Lys Gly
100 105 110

Glu Tyr Leu Asn Arg Asn Ser Phe Val Gln Arg Ser Tyr Ile Arg Gly
115 120 125

Cys Lys Gly Lys Arg Ser Thr His Thr Trp Ile Cys Glu Asn Lys Gly
130 135 140

Asn Asn Asn Ile Cys Ile Pro Asp Arg Arg Val Gln Leu Cys Ile Thr
145 150 155 160

Ala Leu Gln Asp Leu Lys Asn Ser Gly Ser Glu Thr Thr Asp Arg Lys
165 170 175

Leu Leu Arg Asp Lys Val Phe Asp Ser Ala Met Tyr Glu Thr Asp Leu
180 185 190

Leu Trp Asn Lys Tyr Gly Phe Arg Gly Phe Asp Asp Phe Cys Asp Asp
195 200 205

Val Lys Asn Ser Tyr Leu Asp Tyr Lys Asp Val Ile Phe Gly Thr Asp
210 215 220

Leu Asp Lys Asn Asn Ile Ser Lys Leu Val Glu Glu Ser Leu Lys Arg
225 230 235 240

Phe Phe Lys Lys Asp Ser Ser Val Leu Asn Pro Thr Ala Trp Trp Arg
245 250 255

Arg Tyr Gly Thr Arg Leu Trp Lys Thr Met Ile Gln Pro Tyr Ala His

260 265 270

Leu Gly Cys Arg Lys Pro Asp Glu Asn Glu Pro Gln Ile Asn Arg Trp
275 280 285

Ile Leu Glu Trp Gly Lys Tyr Asn Cys Arg Leu Met Lys Glu Lys Glu
290 295 300

Lys Leu Leu Thr Gly Glu Cys Ser Val Asn Arg Lys Lys Ser Asp Cys
305 310 315 320

Ser Thr Gly Cys Asn Asn Glu Cys Tyr Thr Tyr Arg Ser Leu Ile Asn
325 330 335

Arg Gln Arg Tyr Glu Val Ser Ile Leu Gly Lys Lys Tyr Ile Lys Val
340 345 350

Val Arg Tyr Thr Ile Phe Arg Arg Lys Ile Val Gln Pro Asp Asn Ala
355 360 365

Leu Asp Phe Leu Lys Leu Asn Cys Ser Glu Cys Lys Asp Ile Asp Phe
370 375 380

Lys Pro Phe Phe Glu Phe Glu Tyr Gly Lys Tyr Glu Glu Lys Cys Met
385 390 395 400

Cys Gln Ser Tyr Ile Asp Leu Lys Ile Gln Phe Lys Asn Asn Asp Ile
405 410 415

Cys Ser Phe Asn Ala Gln Thr Asp Thr Val Ser Ser Asp Lys Arg Phe
420 425 430

Cys Leu Glu Lys Lys Glu Phe Lys Pro Trp Lys Cys Asp Lys Asn Ser
435 440 445

Phe Glu Thr Val His His Lys Gly Val Cys Val Ser Pro Arg Arg Gln
450 455 460

Gly Phe Cys Leu Gly Asn Leu Asn Tyr Leu Leu Asn Asp Asp Ile Tyr
465 470 475 480

Asn Val His Asn Ser Gln Leu Leu Ile Glu Ile Ile Met Ala Ser Lys
485 490 495

Gln Glu Gly Lys Leu Leu Trp Lys Lys His Gly Thr Ile Leu Asp Asn
500 505 510

Gln Asn Ala Cys Lys Tyr Ile Asn Asp Ser Tyr Val Asp Tyr Lys Asp
515 520 525

Ile Val Ile Gly Asn Asp Leu Trp Asn Asp Asn Asn Ser Ile Lys Val
530 535 540

Gln Asn Asn Leu Asn Leu Ile Phe Glu Arg Asn Phe Gly Tyr Lys Val
545 550 555 560

Gly Arg Asn Lys Leu Phe Lys Thr Ile Lys Glu Leu Lys Asn Val Trp
565 570 575

Trp Ile Leu Asn Arg Asn Lys Val Trp Glu Ser Met Arg Cys Gly Ile
580 585 590

Asp Glu Val Asp Gln Arg Arg Lys Thr Cys Glu Arg Ile Asp Glu Leu
595 600 605

Glu Asn Met Pro Gln Phe Phe Arg Trp Phe Ser Gln Trp Ala His Phe
610 615 620

Phe Cys Lys Glu Lys Glu Tyr Trp Glu Leu Lys Leu Asn Asp Lys Cys

625	630	635	640
Thr Gly Asn Asn Gly Lys Ser Leu Cys Gln Asp Lys Thr Cys Gln Asn			
	645	650	655
Val Cys Thr Asn Met Asn Tyr Trp Thr Tyr Thr Arg Lys Leu Ala Tyr			
	660	665	670
Glu Ile Gln Ser Val Lys Tyr Asp Lys Asp Arg Lys Leu Phe Ser Leu			
	675	680	685
Ala Lys Asp Lys Asn Val Thr Thr Phe Leu Lys Glu Asn Ala Lys Asn			
	690	695	700
Cys Ser Asn Ile Asp Phe Thr Lys Ile Phe Asp Gln Leu Asp Lys Leu			
705	710	715	720
Phe Lys Glu Arg Cys Ser Cys Met Asp Thr Gln Val Leu Glu Val Lys			
	725	730	735
Asn Lys Glu Met Leu Ser Ile Asp Ser Asn Ser Glu Asp Ala Thr Asp			
	740	745	750
Ile Ser Glu Lys Asn Gly Glu Glu Glu Leu Tyr Val Asn His Asn Ser			
	755	760	765
Val Ser Val Ala Ser Gly Asn Lys Glu Ile Glu Lys Ser Lys Asp Glu			
	770	775	780
Lys Gln Pro Glu Lys Glu Ala Lys Gln Thr Asn Gly Thr Leu Thr Val			
785	790	795	800
Arg Thr Asp Lys Asp Ser Asp Arg Asn Lys Gly Lys Asp Thr Ala Thr			
	805	810	815

Asp Thr Lys Asn Ser Pro Glu Asn Leu Lys Val Gln Glu His Gly Thr
820 825 830

Asn Gly Glu Thr Ile Lys Glu Glu Pro Pro Lys Leu Pro Glu Ser Ser
835 840 845

Glu Thr Leu Gln Ser Gln Glu Gln Leu Glu Ala Glu Ala Gln Lys Gln
850 855 860

Lys Gln Glu Glu Glu Pro Lys Lys Lys Gln Glu Glu Glu Pro Lys Lys
865 870 875 880

Lys Gln Glu Glu Glu Gln Lys Arg Glu Gln Glu Gln Lys Gln Glu Gln
885 890 895

Glu Glu Glu Glu Gln Lys Gln Glu Glu Glu Gln Gln Ile Gln Asp Gln
900 905 910

Ser Gln Ser Gly Leu Asp Gln Ser Ser Lys Val Gly Val Ala Ser Glu
915 920 925

Gln Asn Glu Ile Ser Ser Gly Gln Glu Gln Asn Val Lys Ser Ser Ser
930 935 940

Pro Glu Val Val Pro Gln Glu Thr Thr Ser Glu Asn Gly Ser Ser Gln
945 950 955 960

Asp Thr Lys Ile Ser Ser Thr Glu Pro Asn Glu Asn Ser Val Val Asp
965 970 975

Arg Ala Thr Asp Ser Met Asn Leu Asp Pro Glu Lys Val His Asn Glu
980 985 990

Asn Met Ser Asp Pro Asn Thr Asn Thr Glu Pro Asp Ala Ser Leu Lys

995	1000	1005
Asp Asp Lys Lys Glu Val Asp Asp Ala Lys Lys Glu Leu Gln Ser		
1010	1015	1020
Thr Val Ser Arg Ile Glu Ser Asn Glu Gln Asp Val Gln Ser Thr		
1025	1030	1035
Pro Pro Glu Asp Thr Pro Thr Val Glu Gly Lys Val Gly Asp Lys		
1040	1045	1050
Ala Glu Met Leu Thr Ser Pro His Ala Thr Asp Asn Ser Glu Ser		
1055	1060	1065
Glu Ser Gly Leu Asn Pro Thr Asp Asp Ile Lys Thr Thr Asp Gly		
1070	1075	1080
Val Val Lys Glu Gln Glu Ile Leu Gly Gly Gly Glu Ser Ala Thr		
1085	1090	1095
Glu Thr Ser Lys Ser Asn Leu Glu Lys Pro Lys Asp Val Glu Pro		
1100	1105	1110
Ser His Glu Ile Ser Glu Pro Val Leu Ser Gly Thr Thr Gly Lys		
1115	1120	1125
Glu Glu Ser Glu Leu Leu Lys Ser Lys Ser Ile Glu Thr Lys Gly		
1130	1135	1140
Glu Thr Asp Pro Arg Ser Asn Asp Gln Glu Asp Ala Thr Asp Asp		
1145	1150	1155
Val Val Glu Asn Ser Arg Asp Asp Asn Asn Ser Leu Ser Asn Ser		
1160	1165	1170

Val Asp Asn Gln Ser Asn Val Leu Asn Arg Glu Asp Pro Ile Ala
1175 1180 1185

Ser Glu Thr Glu Val Val Ser Glu Pro Glu Asp Ser Ser Arg Ile
1190 1195 1200

Met Thr Thr Glu Val Pro Ser Thr Thr Val Lys Pro Pro Asp Glu
1205 1210 1215

Lys Arg Ser Glu Glu Val Gly Glu Lys Glu Ala Lys Glu Ile Lys
1220 1225 1230

Val Glu Pro Val Val Pro Arg Ala Ile Gly Glu Pro Met Glu Asn
1235 1240 1245

Ser Val Ser Val Gln Ser Pro Pro Asn Val Glu Asp Val Glu Lys
1250 1255 1260

Glu Thr Leu Ile Ser Glu Asn Asn Gly Leu His Asn Asp Thr His
1265 1270 1275

Arg Gly Asn Ile Ser Glu Lys Asp Leu Ile Asp Ile His Leu Leu
1280 1285 1290

Arg Asn Glu Ala Gly Ser Thr Ile Leu Asp Asp Ser Arg Arg Asn
1295 1300 1305

Gly Glu Met Thr Glu Gly Ser Glu Ser Asp Val Gly Glu Leu Gln
1310 1315 1320

Glu His Asn Phe Ser Thr Gln Gln Lys Asp Glu Lys Asp Phe Asp
1325 1330 1335

Gln Ile Ala Ser Asp Arg Glu Lys Glu Glu Ile Gln Lys Leu Leu

1340	1345	1350
Asn Ile Gly His Glu Glu Asp	Glu Asp Val Leu Lys	Met Asp Arg
1355	1360	1365
Thr Glu Asp Ser Met Ser Asp	Gly Val Asn Ser His	Leu Tyr Tyr
1370	1375	1380
Asn Asn Leu Ser Ser Glu Glu	Lys Met Glu Gln Tyr	Asn Asn Arg
1385	1390	1395
Asp Ala Ser Lys Asp Arg Glu	Glu Ile Leu Asn Arg	Ser Asn Thr
1400	1405	1410
Asn Thr Cys Ser Asn Glu His	Ser Leu Lys Tyr Cys	Gln Tyr Met
1415	1420	1425
Glu Arg Asn Lys Asp Leu Leu	Glu Thr Cys Ser Glu	Asp Lys Arg
1430	1435	1440
Leu His Leu Cys Cys Glu Ile	Ser Asp Tyr Cys Leu	Lys Phe Phe
1445	1450	1455
Asn Pro Lys Ser Ile Glu Tyr	Phe Asp Cys Thr Gln	Lys Glu Phe
1460	1465	1470
Asp Asp Pro Thr Tyr Asn Cys	Phe Arg Lys Gln Arg	Phe Thr Ser
1475	1480	1485
Met His Tyr Ile Ala Gly Gly	Gly Ile Ile Ala Leu	Leu Leu Phe
1490	1495	1500
Ile Leu Gly Ser Ala Ser Tyr	Arg Lys Asn Leu Asp	Asp Glu Lys
1505	1510	1515

Gly Phe Tyr Asp Ser Asn Leu Asn Asp Ser Ala Phe Glu Tyr Asn
1520 1525 1530

Asn Asn Lys Tyr Asn Lys Leu Pro Tyr Met Val Val Asp Gln Gln
1535 1540 1545

Ile Asn Val Val Asn Ser Asp Leu Tyr Ser Glu Gly Ile Tyr Asp
1550 1555 1560

Asp Thr Thr Thr Phe
1565

<210> 13

<211> 24

<212> DNA

<213> artificial sequence

<220>

<223> sequence is completely synthesized

<400> 13
cctggagccc gtcagtatcg gcgg

24

<210> 14

<211> 23

<212> DNA

<213> artificial sequence

<220>

<223> sequence is completely synthesized

<400> 14

ggtagcgacc ggcgctcagc tgg

23

<210> 15

<211> 25

<212> DNA

<213> artificial sequence

<220>

<223> sequence is completely synthesized

<400> 15

aaaagtgatg atagaaatgc ttgtg

25

<210> 16

<211> 25

<212> DNA

<213> artificial sequence

<220>

<223> sequence is completely synthesized

<400> 16

ttttgtgat cttacttatt tcacc

25

<210> 17

<211> 22

<212> DNA

<213> artificial sequence

<220>

<223> sequence is completely synthesized

<400> 17

cggaatcagg tttaaatacca ac

22

<210> 18

<211> 21

<212> DNA

<213> artificial sequence

<220>

<223> sequence is completely synthesized

<400> 18

agatcgtttt tcatcagggg g

21